STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/587.324
Source:	IFWP.
Date Processed by STIC:	08/04/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR	DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/587, 324
ATTN:	NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
1	_Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	_Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3	_Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.
4	_Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5	_Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	_PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7	_Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8	_Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10	_Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)
11/	_Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown. Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	_ Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/587,324

DATE: 08/04/2006

TIME: 14:30:53

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\08042006\J587324.raw

```
3 <110 > APPLICANT: Geerlings, Arjian
               Hidalgo Zarco, Fernando
               Boza Puerta, Julio
               Jimenez Lopez, Jesus
      8 <120> TITLE OF INVENTION: ANTIHYPERTENSIVE PEPTIDES
     10 <130> FILE REFERENCE: 4258-122
                                                                    Does Not Comply
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/587,324
                                                                    Corrected Diskette Needed
     13 <141> CURRENT FILING DATE: 2006-07-26
     15 <150> PRIOR APPLICATION NUMBER: PCT/EP2005/001992
     16 <151> PRIOR FILING DATE: 2005-02-23
     18 <150> PRIOR APPLICATION NUMBER: EP04075587.8
     19 <151> PRIOR FILING DATE: 2004-02-26
     21 <160> NUMBER OF SEQ ID NOS: 9
                                                           L2137 Response is s
Attricial, Pls Explains
Source of genetic
Haleval. See Glem
Maleval. See Glem
11 on zeroy Sheet.
78 cmm 2007 Sheet.
     23 <170> SOFTWARE: PatentIn version 3.3
     25 <210> SEQ ID NO: 1
     26 <211> LENGTH: 6
     27 <212> TYPE: PRT
     28 <213> ORGANISM: Artificial
     30 <220> FEATURE:
     31 <223> OTHER INFORMATION \ Artificial
     33 <400> SEQUENCE: 1
     35 Tyr Gly Pro Ile Pro Asn
     36 1
                          5
     39 <210> SEQ ID NO: 2
     40 <211> LENGTH: 4
     41 <212> TYPE: PRT
     42 <213> ORGANISM: Artificial
     44 <220> FEATURE:
     45 <223 > OTHER INFORMATION: (Artificial
     47 <400> SEQUENCE: 2
     49 Ser Leu Pro Glu
     50 1
     53 <210> SEQ ID NO: 3
     54 <211> LENGTH: 4
     55 <212> TYPE: PRT
     56 <213> ORGANISM: Artificial
     58 <220> FEATURE:
     59 <223 > OTHER INFORMATION: Artificial
     61 <400> SEQUENCE: 3
     63 Ser Glu Pro Lys
     64 1
     67 <210> SEQ ID NO: 4
     68 <211> LENGTH: 8
```

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/587,324**DATE: 08/04/2006

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```
69 <212> TYPE: PRT
70 <213> ORGANISM: Artificial
72 <220> FEATURE:
73 <223 > OTHER INFORMATION \ Artificial
75 <400> SEQUENCE: 4
77 Phe Tyr Gly Pro Ile Pro Asn Ser
78 1
81 <210> SEQ ID NO: 5
82 <211> LENGTH: 6
83 <212> TYPE: PRT
84 <213> ORGANISM: Artificial
86 <220> FEATURE:
87 <223> OTHER INFORMATION Artificial
89 <400> SEQUENCE: 5
91 Asn Ser Leu Pro Glu Asn
92 1
95 <210> SEQ ID NO: 6
96 <211> LENGTH: 6
97 <212> TYPE: PRT
98 <213> ORGANISM: Artificial
100 <220> FEATURE:
101 <223 > OTHER INFORMATION Artificial
103 <400> SEQUENCE: 6
105 Leu Ser Glu Pro Lys Val
106 1
109 <210> SEQ ID NO: 7
110 <211> LENGTH: 5
111 <212> TYPE: PRT
112 <213> ORGANISM: Artificial
114 <220> FEATURE:
115 <223> OTHER INFORMATION Artificial
117 <400> SEQUENCE: 7
119 Gly Pro Ile Pro Asn
120 1
123 <210> SEQ ID NO: 8
124 <211> LENGTH: 3
125 <212> TYPE: PRT
126 <213> ORGANISM: Artificial
128 <220> FEATURE:
129 <223 > OTHER INFORMATION: (Artificial
131 <400> SEQUENCE: 8
133 Leu Pro Glu
134 1
137 <210> SEQ ID NO: 9
138 <211> LENGTH: 3
139 <212> TYPE: PRT
140 <213> ORGANISM: Artificial
142 <220> FEATURE:
143 <223> OTHER INFORMATION: Artificial
```

Same Error

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Input Set : A:\PTO.KD.txt

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145 <400> SEQUENCE: 9

147 Glu Pro Lys

148 1

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/587,324

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Input Set : A:\PTO.KD.txt

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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9

VERIFICATION SUMMARY

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Input Set : A:\PTO.KD.txt

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L:12 M:270 C: Current Application Number differs, Replaced Current Application

Number